

# Package: dam (via r-universe)

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**Title** Data Analysis Metabolomics

**Version** 0.0.1

**Description** A collection of functions which aim to assist common computational workflow for analysis of matabolomic data..

**Depends** R (>= 3.2.3)

**License** CC0

**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 5.0.1.9000

**NeedsCompilation** no

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**Repository** <https://slakov.r-universe.dev>

**RemoteUrl** <https://github.com/cran/dam>

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correctedMarkers      *A correctedMarkers Function*

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**Description**

This function allows you to perform several multiple regression on metabolomic data.

**Usage**

```
correctedMarkers(dat, fac, con, comp)
```

**Arguments**

dat	Data matrix.
fac	Index of the group of samples.
con	Index of correction variable.
comp	Index (range) of measured compounds.

**Examples**

```
correctedMarkers(iris, 5, 1, 2:4)
```

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